Amendment to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1. (currently amended) A method of manufacturing a glycoprotein having a mammalian-type or human-type sugar chain, comprising a step in which a transformed plant cell is [obtained] produced by introducing to a plant cell [the] a gene encoding a [of] glycosyltransferase enzyme and [the] a gene of an exogenous glycoprotein, and a step in which the [obtained] produced transformed plant cell is cultivated.

2. (currently amended) The method according to claim 1, wherein the glycosyltransferase enzyme is an enzyme [capable of conducting a transfer reaction of] which transfers a galactose residue to a non-reducing terminal acetylglucosamine residue.

3. (currently amended) The method according to claim 1, wherein the glycoprotein with a mammalian-type or human-type sugar chain comprises a core sugar chain and an outer sugar chain, wherein the core sugar chain comprises a plurality of mannose and acetylglucosamine, and wherein the outer sugar chain contains a terminal sugar chain portion with a non-reducing terminal galactose.

4. (previously presented) The method according to claim 3, wherein the outer sugar chain has a straight chain configuration.

5. (previously presented) The method according to claim 3, wherein the outer sugar chain has a branched configuration.

6. (currently amended) The method according to claim 5, wherein the branched sugar chain portion has a mono-, bi-, tri-, or tetra-configuration.

7. (currently amended) The method according to claim 1, wherein the glycoprotein contains neither fucose nor xylose linked to one or more of the core sugar chain, the outer sugar chain and the terminal sugar chain.

8. (currently amended) A transformed plant cell having a sugar chain adding mechanism which can conduct a transfer reaction of a galactose residue to a non-reducing terminal acetylglucosamine residue, wherein the sugar chain adding mechanism adds a sugar chain containing a core sugar chain and an outer sugar chain, wherein the core sugar chain comprises a plurality of mannose and acetylglucosamine, and wherein the outer sugar chain contains a terminal sugar chain portion with a non-reducing terminal galactose.

9. (currently amended) The transformed plant cell according to claim 8, wherein the plant cell is transformed with the gene of a first enzyme [capable of conducting a transfer reaction of] which transfers a galactose residue to a non-reducing terminal acetylglucosamine residue and the gene of a second enzyme which can [enhance] improve the performance of the first enzyme.

10. (currently amended) [A] The transformed plant cell according to claim 9, wherein the second enzyme is selected from the group consisting of Mannosidase I, Mannosidase II, [β1,4-Glactosyltransferase (GalT)] and N-acetylglycosaminyltransferase I (GlcNAc I).

11. (currently amended) A transformed plant regenerated from the plant cell of claim 8.

12. (currently amended) A recombinant plant, or portion thereof, that produces mammalian-type glycoproteins comprising neither fucose or xylose linked to one or more of the core sugar chain, the outer sugar chain and the terminal sugar chain.

13. (cancelled)

14. (cancelled)

15. (new) A method of manufacturing a glycoprotein, which method includes introducing into a plant cell a gene encoding a glycosyltransferase enzyme of human origin selected from the group consisting of galactosyltransferase, galactosidase and β-galactosidase and a gene encoding an exogenous glycoprotein selected from one or more of the group consisting of enzymes, hormones, cytokines, antibodies, vaccines, receptors and serum proteins, the

glycoprotein produced including a core sugar chain including a plurality of mannose and acetylglucosamine residues, and an outer sugar chain containing a terminal sugar chain portion with a non-reducing terminal galactose, and wherein the glycoprotein produced has no fucose or xylose linked to one or more of the core sugar chain, the outer sugar chain and the terminal sugar chain.

16. (new) The method according to claim 15, which method further includes introducing a gene encoding a second enzyme capable of enhancing the efficiency of the glycosyl transferase enzyme.

17. (new) The method according to claim 16, wherein the second enzyme is selected from the group consisting of Mannosidase I, Mannosidase II, and N-acetylglycosaminyltransferase I (GlcNAc I).

18. (new) The method according to claim 15, wherein the exogenous glycoprotein encoded by the introduced gene is an enzyme selected from one or more of the group consisting of horseradish peroxidase, kinase, glucocerebrosidase, α-galactosidase, tissue-type plasminogen activator (TPA), and 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase.

19. (new) The method according to claim 15, wherein the exogenous glycoprotein encoded by the introduced gene is a hormone or cytokine selected from one or more of the group consisting of enkephalin, interferon-alpha, granulocyte-macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), chorion stimulating hormone, interleukin-2, interferon-beta, interferon-gamma, erythropoietin, vascular endothelial growth factor, human choriogonadotropin (HCG), leuteinizing hormone (LH), thyroid stimulating hormone (TSH), prolactin, and ovary stimulating hormone.

20. (new) The method according to claim 15, wherein the exogenous glycoprotein encoded by the introduced gene is an antibody selected from immunoglobulin G (IgG) or single chain variable region antibody fragments (scFV).

21. (new) The method according to claim 15, wherein the plant cell is derived from a plant selected from the group consisting of plants in the families of

Solanaceae, Poaeae, Brassicaceae, Rosaceae, Leguminosae, Curcurbitaceae, Lamiaceae, Liliaceae, Chenopodiaceae, and *Umbelliferae*.

22. (new) The method according to claim 21, wherein the plant cell is derived from a plant selected from the group consisting of tobacco, tomato, potato, rice, maize, radish, soybean, peas, alfalfa, or spinach.

23. (new) The method according to claim 1, 2 or 3 wherein the sugar chain is a mammalian-type sugar chain.

24. (new) The method according to claim 1, 2 or 3 wherein the sugar chain is a human-type sugar chain.